

(A) a polypeptide which is capable of inducing cell death (apoptosis) and comprises the amino acid sequence of SEQ ID NO:2;

(B) a polypeptide which has a property of being capable of inducing cell death and has at least 85% sequence identity to the amino acid sequence of SEQ ID NO:2;

B¹
cont. (C) a fragment of a polypeptide of SEQ ID NO:2 which is capable of inducing cell death;

(D) a fragment which is capable of inducing cell death and has at least 85% sequence identity to fragment (C);

(E) a fragment of a polypeptide of SEQ ID NO:2 which lacks the property of being capable of inducing cell death and which inhibits the ability of the polypeptide (A) or (B) to induce cell death; and

(F) a fragment which lacks the property of being capable of inducing cell death and which inhibits the ability of the polypeptide (A) or (B) to induce cell death, said fragment having at least 85% sequence identity to fragment (E).

B²
8 (Amended). A polypeptide capable of inducing cell death, consisting of an amino acid sequence selected from the group consisting of amino acid residues 13 to 275 of SEQ ID NO:2 and an amino acid sequence having at least 85% sequence identity to residues 13 to 275 of SEQ ID NO:2.

B3
13(Amended). A polypeptide capable of inhibiting the ability of the polypeptide of SEQ ID NO:2 to induce cell death, consisting of an amino acid sequence selected from the group consisting of amino acid residues 321 to 360 of SEQ ID NO:2 and an amino acid sequence having at least 85% sequence identity to residues 321 to 360 of SEQ ID NO:2.

Please add new claims 31-36 as follows:

--31(New). The polypeptide of claim 1, wherein said polypeptide (B) or said fragment (D) has at least 90% sequence identity to the amino acid sequence of SEQ ID NO:2 or to the fragment of (C), respectively.--

--32(New). The polypeptide of claim 1, wherein said polypeptide (B) or said fragment (D) has at least 95% sequence identity to the amino acid sequence of SEQ ID NO:2 or to the fragment of (C), respectively.--

--33(New). The polypeptide of claim 8 which has at least 90% sequence identity to residues 13 to 275 of SEQ ID NO:2.--

--34(New). The polypeptide of claim 8 which has at least 95% sequence identity to residues 13 to 275 of SEQ ID NO:2.--

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--35(New). The polypeptide of claim 13 which has at least 90% sequence identity to residues 321 to 360 of SEQ ID NO:2.--

--36(New). The polypeptide of claim 13 which has at least 95% sequence identity to residues 321 to 360 of SEQ ID NO:2.--
